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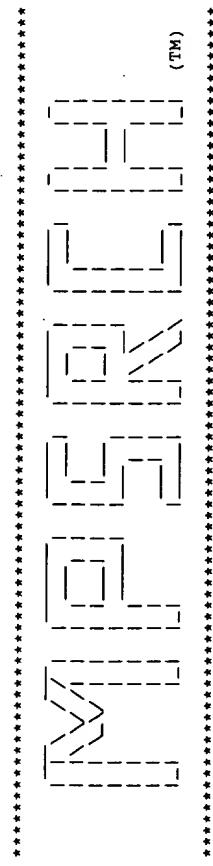
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MPsrch_PP protein - protein database search, using Smith-Waterman algorithm

Run on: Mon Oct 2 09:33:11 2000: MasPar time 6.68 Seconds
 436.063 Million cell updates/sec

Tabular output not generated.

Title: >US-09-381-497-2
 Description: (1-123) from US09381497.pep
 Perfect Score: 903
 Sequence: 1 EVQLYVEGGGLVKGPGSSKLK.....SSYGYLFAYKGQGTLYVSA 123

Scoring table: PAM 150

Gap 11

Searched: 188963 seqs, 23686106 residues

Post-processing: Minimum Match 0%

Listing first 45 summaries

Database: a-geneseq36

1:geneseq36

Statistics: Mean 30.324; Variance 152.652; scale 0.199

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description	Pred. No.
1	896	99.2	123	1 W66099	anti-CD22 monoclonal a	1.58e-63
2	748	82.8	121	1 W86125	Protein sequence of mo	3.76e-51
3	744	82.4	121	1 W22951	Monoclonal antibody (M	8.11e-51
4	744	82.4	121	1 W86118	Murine 340 Vh amino ac	8.11e-51
5	735	81.4	139	1 W21656	Chimeric MAb 15 PCR mo	4.56e-50
6	730	80.8	140	1 W21654	Mouse MAb 15 heavy cha	1.19e-49
7	728	80.6	136	1 R06251	Variable region of mur	1.75e-49
8	717	79.4	139	1 R52773	Murine KC-4 immunoglob	1.44e-48
9	716	79.4	139	1 R52791	Murine KC-4 immunoglob	1.44e-48
10	712	79.3	118	1 W89627	Mouse humanised antibo	1.74e-48
11	716	79.3	118	1 W57576	Chimeric H Chain SEQ I	1.74e-48
12	716	79.3	137	1 W89625	Mouse humanised antibo	1.74e-48
13	716	79.3	137	1 W57592	Chimeric antibody agai	1.74e-48
14	716	79.3	247	1 W11917	Murine MAb SK48-E26 he	1.74e-48
15	713	79.0	121	1 W86122	Protein sequence of de	3.10e-48
16	713	79.0	123	1 W08582	Human antibody C4.1 he	3.10e-48
17	712	78.8	121	1 W86120	Protein sequence of hu	3.75e-48
18	712	78.8	139	1 W21652	Humanised reshaped MAb	3.75e-48
19	710	78.6	138	1 R20064	MRK16-H chain.	5.50e-48
20	709	78.5	117	1 W76003	Lm609 antibody heavy c	6.67e-48
21	706	78.2	142	1 R30882	Antibody 4A2 heavy cha	1.18e-47
22	704	78.0	138	1 W03722	Anti-human IgP9 MAb 39	1.74e-47
23	703	77.9	123	1 R43827	Anti-lysozyme VH.	2.10e-47

RESULT
 ID: W66099 standard; Protein: 123 AA.
 AC: W66099.
 DT: 10-DEC-1998 (first entry)
 DE: anti-CD22 monoclonal antibody heavy chain variable region.
 KW: anti-CD22 monoclonal antibody heavy chain variable region; VI;
 Pseudomonas exotoxin; variable heavy chain; VH; variable light chain;
 KW: malignant B-cell; immunodiagnosis; RFB4 IgG.
 OS: Mammalia.
 FH: Key
 FT: Misc_difference 121
 /note= "Encoded by gtc"
 PN: W09841641-A1.
 PD: 24-SEP-1998.
 PF: 19-MAR-1998; U05453
 PR: 20-MAR-1997; US-041437.
 (USSH) US DEPT HEALTH & HUMAN SERVICES.
 PI: Fitzgerald D, Kreitman R, Mansfield E, Pastan I;
 WP1: 98-5212/27/44.
 DR: N-PSDB; V07642.

PT: Recombinant anti-CD22 antibodies and immuno-conjugates - of
 PT: antibodies linked to a therapeutic agent, e.g. Pseudomonas exotoxin
 PT: or a label; for inhibiting malignant B-cells
 PS: Claim 6; Fig 1; 71PP; English.
 CC: The invention claims for a recombinant immunotoxin comprising
 CC: a therapeutic agent (e.g. Pseudomonas exotoxin) or a detectable
 CC: label peptide bonded to a recombinant anti-CD22 antibody (RFB4 IgG)
 CC: having the present variable heavy (VH) chain with a cysteine residue
 CC: at amino acid 44 and a variable light (VL; W66098) chain with a
 CC: cysteine residue at amino acid 100. The immunotoxin is claimed
 CC: to inhibit the growth of malignant B-cells in vivo, such as rodent,
 CC: canine or primate B-cells. The anti-CD22 antibody is claimed useful
 CC: for detecting CD22 protein in a sample or
 CC: can be used in diagnostic kits.
 SQ: Sequence 123 AA;

Query Match 99.2%; Score 896; DB 1; Length 123;
 Best Local Similarity 99.2%; Pred. No. 1.58e-03;
 Matches 122; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
 Db 1 EVOLYESGGGLVKGPGSSKLKSCAASGFATSYDMSWRQTPKLEWAVISSLGGTTYY 60
 1 EVOLYESGGGLVKGPGSSKLKSCAASGFATSYDMSWRQTPKLEWAVISSLGGTTYY 60
 1 PDTVKGRFTISRDNAKNTLYLQMSSLLKSCAASGFATSYDMSWRQTPKLEWAVISSLGGTTYY 60
 1 PDTVKGRFTISRDNAKNTLYLQMSSLLKSCAASGFATSYDMSWRQTPKLEWAVISSLGGTTYY 60
 1 PDTVKGRFTISRDNAKNTLYLQMSSLLKSCAASGFATSYDMSWRQTPKLEWAVISSLGGTTYY 60

Db 61 PDTVKGRFTISRDNAKNTLYLQMSSLLKSCAASGFATSYDMSWRQTPKLEWAVISSLGGTTYY 120
 1 EVOLYESGGGLVKGPGSSKLKSCAASGFATSYDMSWRQTPKLEWAVISSLGGTTYY 120
 1 PDTVKGRFTISRDNAKNTLYLQMSSLLKSCAASGFATSYDMSWRQTPKLEWAVISSLGGTTYY 120
 1 PDTVKGRFTISRDNAKNTLYLQMSSLLKSCAASGFATSYDMSWRQTPKLEWAVISSLGGTTYY 120

RESULT	2	standard; Protein; 121 AA.	31..35 /note= "Complementarity-determining region 1" 36..55 /note= "Framework 2" 56..56 /note= "Complementarity-determining region 2" 67..98 /note= "Framework 3" 97..110 /note= "Complementarity-determining region 3" 111..121 /note= "Framework 4"
Db	121	TS4 123	FT Region
AC	W86125;	1	FT Region
Qy	121	VSA 123	FT Region
RESULT	2	standard; Protein; 121 AA.	FT Region
AC	W86125;		FT Region
DT	03-MAR-1999	(first entry)	FT Region
DE	Protein sequence of mouse de-immunised 340 Vh.	immune system; SK;	FT Region
KW	Non-immunogenic; epitope; T-cell; immunogenicity;	immunogen; murine.	FT Region
KW	Immunoglobulin; therapeutic; streptokinase; de-immunised; murine.	OS	FT Region
OS	Mus sp.	OS	FT Region
PN	W09852976-A1.	PN W09730354-A2.	PN W09730354-A2.
PR	26-NOV-1998.	PD 21-AUG-1997	PD 21-AUG-1997
PF	21-MAY-1998; GB-01473.	PF 17-FEB-1997	PF 17-FEB-1997
PR	14-APR-1998; GB-007751.	PR 16-FEB-1996	PR 16-FEB-1996
PR	21-MAY-1997; GB-010480.	PA (UNIV NOTTINGHAM.	PA (UNIV NOTTINGHAM.
PR	31-JUL-1997; GB-016197.	PI Durrant LG, Liu DT;	PI Durrant LG, Liu DT;
PR	28-NOV-1997; GB-025270.	DR WPI: 97-425169/39.	DR WPI: 97-425169/39.
PR	02-DEC-1997; US-067235.	PR N-PSDB; T75983.	PR N-PSDB; T75983.
PA	(BIOV-) BIOVATION LTD.	PT Foetal cell analysis comprising isolating at least 2 sorts of foetal	PT Foetal cell analysis comprising isolating at least 2 sorts of foetal
PT	Carr EU;	cell from maternal sample - specifically nucleated red cells and	cell from maternal sample - specifically nucleated red cells and
PT	WPI; 99-045301/04.	PT trophoblasts, useful for pre-natal diagnosis of genetic	PT trophoblasts, useful for pre-natal diagnosis of genetic
DR	Reducing immunogenicity of proteins - by modifying the amino acid	PT abnormalities or foetal sex.	PT abnormalities or foetal sex.
PT	sequence of the protein to eliminate potential epitopes for T-cells	PS Example 2; Page 15; 31pp; English.	PS Example 2; Page 15; 31pp; English.
PT	of a given species	CC This is the heavy chain variable region of a monoclonal antibody (MAB)	CC This is the heavy chain variable region of a monoclonal antibody (MAB)
PT	Example 2; Fig 10; 77pp; English.	CC 340 acts as a trophoblast binding agent in a method for	CC 340 acts as a trophoblast binding agent in a method for
PS	CC The invention relates to a method for the production of non-immunogenic	CC analysing human foetal cells. The method comprises isolating at least 2	CC analysing human foetal cells. The method comprises isolating at least 2
CC	CC proteins. The method comprises determining at least part of the amino	CC types of nucleated fetal cell from a maternal sample. The foetal cells	CC types of nucleated fetal cell from a maternal sample. The foetal cells
CC	CC acid sequence of the protein; (b) identifying in the amino acid sequence	CC are nucleated red cells (erythrocytes) and trophoblasts.	CC are nucleated red cells (erythrocytes) and trophoblasts.
CC	CC one or more potential epitopes for T-cells (T-cell epitopes) of the given	CC The trophoblasts are isolated by contacting the maternal sample with a	CC The trophoblasts are isolated by contacting the maternal sample with a
CC	CC species; and (c) modifying the amino acid sequence to eliminate at least	CC epidermal growth factor receptor (EGFR). MAB340 comprising at	CC epidermal growth factor receptor (EGFR). MAB340 comprising at
CC	CC one of the T-cell epitopes identified in step (b) thereby to eliminate or	CC least the CDR regions of one or other of the light or heavy chain	CC least the CDR regions of one or other of the light or heavy chain
CC	CC reduce the immunogenicity of the protein when exposed to the immune	CC variable regions of MAB 340 is used as the binding agent. Erythrocytes	CC variable regions of MAB 340 is used as the binding agent. Erythrocytes
CC	CC system of the given species. A method of analysing a pre-existing protein	CC are also isolated with a specific anti-transferrin antibody. The target	CC are also isolated with a specific anti-transferrin antibody. The target
CC	CC to predict the basis for immunogenicity is also provided. The	CC nucleotide sequences in the foetal cells are amplified by PCR primers.	CC nucleotide sequences in the foetal cells are amplified by PCR primers.
CC	CC methods can be used particularly for reducing the immunogenicity of	CC The isolated cells can be used for genetic and biochemical analysis,	CC The isolated cells can be used for genetic and biochemical analysis,
CC	CC immunoglobulins or therapeutic proteins, e.g. Streptokinase (SK). The	CC particularly to determine the sex of the foetus or to detect inherited	CC particularly to determine the sex of the foetus or to detect inherited
CC	CC products can be used for diagnosis and therapy. The present sequence	CC abnormalities, e.g. Down's syndrome. More generally MAB 340 can be used	CC abnormalities, e.g. Down's syndrome. More generally MAB 340 can be used
CC	CC represents the protein sequence of mouse de-immunised 340 Vh.	CC to enrich (for subsequent analysis) rare malignant cells, e.g. squamous	CC to enrich (for subsequent analysis) rare malignant cells, e.g. squamous
CC	CC sequence 121 AA;	CC carcinoma cells, that express EGFR receptor. The method is not	CC carcinoma cells, that express EGFR receptor. The method is not
CC	CC significantly invasive and does not carry the risk of inducing abortion.	CC significantly invasive and does not carry the risk of inducing abortion.	CC significantly invasive and does not carry the risk of inducing abortion.
CC	CC use of two types of cells significantly improves the sensitivity of	CC use of two types of cells significantly improves the sensitivity of	CC use of two types of cells significantly improves the sensitivity of
CC	CC pre-natal diagnosis.	CC pre-natal diagnosis.	CC pre-natal diagnosis.
SO	SO Sequence 121 AA;	SO Sequence 121 AA;	SO Sequence 121 AA;
Query Match	82.8%	Score 744; DB 1; Length 121;	Score 744; DB 1; Length 121;
Best Local Similarity	87.8%	Pred. No. 3.76e-51;	Best Local Similarity 87.7%; Pred. No. 9.11e-51;
Matches	103;	Indels 2;	Matches 103; Indels 2;
Conservative	7;	Matches 11;	Conservative 7; Mismatches 11;
Query Match	82.4%	Score 744; DB 1; Length 121;	Score 744; DB 1; Length 121;
Db	1	EVOLVESGGGLYKAGGSIKLSCAASGPAFDYDMSWROTPEKRLEAVYIGGGDFTYY 60	Best Local Similarity 87.7%; Pred. No. 9.11e-51;
Qy	1	EVOLVESGGGLYKPGGSIKLSCAASGRAFSIDMSWROTPEKRLEAVYISGGGTTYY 60	Matches 103; Conservative 7; Mismatches 11;
Db	61	PDTVKGRFTISRDNGKNSLYLQMLNSLKSEDTAMYCARHYVHDYAV-DYWGOGSTVT 118	Db 1 EVOLVESGGGLYKPGGSIKLSCAASGRAFSIDMSWROTPEKRLEAVYIGGGDFTYY 60
Qy	61	PDTVKGRFTISRDNAKNTLYLQMLSSLKSEDTAMYCARHYGSSYCVLFAYWQGTLVT 120	Qy 1 EVOLVESGGGLYKPGGSIKLSCAASGRAFSIDMSWROTPEKRLEAVYISGGGTTYY 60
Db	119	VSS 121	Db 61 PDTVKGRFTISRDNGKNTLYLQMLNSLKSEDTAMYCARHYVHDYAV-DYWGOGSTVT 118
Qy	121	VSA 123	Qy 61 PDTVKGRFTISRDNAKNTLYLQMLSSLKSEDTAMYCARHYGSSYCVLFAYWQGTLVT 120
Db	119	VSS 121	Db 119 VSS 121
Qy	121	VSA 123	Qy 121 VSA 123
RESULT	3	standard; Protein; 121 AA.	RESULT 4
AC	W22951;		ID W86118 standard; Protein; 121 AA.
DT	02-APR-1998	(first entry)	AC W86118.
DE	Monoclonal antibody (Mab) 340 heavy chain variable region.	DT 03-MAR-1999 (first entry)	
KW	Mab 340; monoclonal antibody; foetal cell; pre-natal diagnosis;	DE Marine 340 Vh amino acid sequence.	
KW	erythrocyte; trophoblast; genetic abnormality; foetal sex; analysis;	KW Non-immunogenic; epitope; T-cell; immune system; SK;	
KW	epidermal growth factor receptor; EGFR; Down's syndrome; human.	KW Immunoglobulin; therapeutic; streptokinase; diagnosis; murine.	
OS	Mus sp.	OS W09852976-A1.	
PH	Key	Location/Qualifiers	Location/Qualifiers
	1..30	note= "Framework 1"	note= "Framework 1"
	Region		

Query	Match	Score	Length	DB
Best Local Matches	82.4%; Similarity 83.7%; Conservative 103;	744; Pred. No. 8.11e-51;	121;	
Qy				
Qy	1 EVQLYESGGGLVKGAGSLKLSCAASGFAFDYDMSMWQTPEKRLLEWVAYIGSGGDRYY 60			
Qy	1 EVQLYESGGGLVKGAGSLKLSCAASGFAFDYDMSMWQTPEKRLLEWVAYIGSGGDRYY 60			
Qy	61 PDTVKRGFTISRDNGKNTLYLQLNSLKSEDTAMYCARHGYHDYAV- -DYGQGTSTV 118			
Qy	61 PDTVKRGFTISRDNGKNTLYLQLNSLKSEDTAMYCARHGYGSYGVLFAWGQPLVT 120			
Qy				
Qy	119 VSS 121			
Qy	121 VSA 123			

RESULT	5
ID	W21656 standard; Protein; 139 AA.
AC	W21656;
DT	03-JAN-1998 (first entry)
DE	Chimeric Mab 15 PCR-modified heavy chain variable region.
KW	Humanised antibody; monoclonal antibody; Mab 15; tumour; lung cancer; therapy; Chimeric Mab musculus.
DS	Chimeric synthetic.
CCS	
PH	
Key	Location/Qualifiers
Peptide	1..19
	/label= Sig_peptide
Protein	25..139
	/label= Mat_protein
Region	20..49
	/label= Framework-1
Region	50..54
	/label= CDR1
	/note= "complementarity determining region
Region	55..68
	/label= Framework-2
Region	69..86
	/label= CDR2
	/note= "complementarity determining region
Region	87..117
	/label= Framework-3

FT	Region	118..128
FT		/label=" CDR3
FT		/note=" Complementarity determining region 3"
FT	Region	129..138
FT		/label=" Framework-4
PN	EP-781847-A1.	
PD	02-JUL-1997.	
PF	25-OCT-1996; 117154	
PR	06-NOV-1995; EP-117407.	
PA	(MERE) MERCK PATENT GMBH.	
PT	Bendig, M., Jones, T.,	
DR	WPI; 97-33904/31.	
DR	N-PSDB; T72269.	
PT	Humanised form of murine monoclonal antibody MAB 15 - useful for	
PT	treating lung cancer	
PS	Disclosure: Fig 5; 71PP; English.	
CC	This polypeptide comprises the heavy chain variable region VH	
CC	region of murine monoclonal antibody (MAB) 15 (DSM ACC2117). It is	
CC	encoded by a 15 VH cDNA sequence (T72269) modified for the	
CC	expression of a chimeric antibody. The VL sequence was similarly	
CC	obtained (see W1655). The modified VH and VL sequences were used	
CC	in a claimed process to model and design novel-humanised,-reshaped	
CC	MAB 15, having humanised, reshaped VH and VL sequences (see W1652	
CC	and W1651), which can be used for treating tumours, especially	
CC	lung cancer, and for the manufacture of a drug related to tumours,	
SQ	sequence 139 AA;	

Best Local Matches		Similarity	Length	1
Db	EVQVYEGGLVKPQGGSLKLSAACAGFAFSYDMSWVRQPKEKRLIEWAYLRSRGGSSTYY	8.9%	50	Pred. No. 4.5e-50;
Qy	EVQVYEGGLVKPQGGSLKLSAACAGFAFSYDMSWVRQPKEKRLIEWAYLRSRGGSSTYY	9.1	50	Matches 10%; conservative
Db	PTDVKRGRETISRDNAKILFLQMTSLKSEDTAMYCARHGEEVPRW	8.9	137	Length 137;
Qy	PTDVKRGRETISRDNAKILFLQMTSLKSEDTAMYCARHGEEVPRW	8.9	137	Matches 10%; conservative
Db	PTDVKRGRETISRDNAKILFLQMTSLKSEDTAMYCARHSGYGSSTYY	9.0	136	Pred. No. 4.5e-50;
Qy	PTDVKRGRETISRDNAKILFLQMTSLKSEDTAMYCARHSGYGSSTYY	9.0	136	Matches 10%; conservative
Db	VSA 1.39	1.39	137	
Qy	VSA 1.33	1.33	121	

ID W21654 standard; Protein; 140 AA.
 AC W21654;
 DT 03-JAN-1998 (first entry)
 DE Mouse MAb 15 heavy chain variable region.
 KW Humanised antibody; monoclonal antibody; Mab 15; tumour;
 KW lung cancer; therapy.
 OS Mus musculus.
 FH Key Location/Qualifiers
 Peptide 1..19
 FT /label= sig_peptide
 Protein 20..140
 FT /label= Mat_protein
 Region 20..49
 FT /label= Framework-1
 Region 50..53
 FT /label= CDR1
 FT /note= "complementarity determining region 1"
 Region 54..68
 FT /label= Framework-2
 Region 69..86
 FT /label= CDR2
 FT /note= "complementarity determining region 2"
 Region 87..117
 FT /label= Framework-3
 Region 118..128
 FT /label= CDR3
 FT

Region	129.	139
FT	FT	/label= Framework-4
EP	EP-781847-A1.	
PN	02-JUL-1997.	
PD	23-OCT-1996.	
PP	117154.	
EP	06-NOV-1995: EP-117407	
PA	(MERE) MERCK PATENT GMBH.	
PI	Bandig M., Jones T., Saldana J;	
WPI	97-334804/31.	
NP	W72267.	
DR	N-PSDB; W72267.	
PR	Humanised form of murine monoclonal antibody MAB 15 - useful for treating lung cancer	
PS	Example 1: Fig 2; 71PP; English.	
CC	This polypeptide comprises the heavy chain variable region VH of murine monoclonal antibody (MAB) 15 (DSM Acc2117), a MAB that shows a therapeutic effect on human tumour cells, especially human lung cancer. Its sequence was deduced from an isolated cDNA cloner (see T72267). The MAB 15 VH region sequence (W71653) has also been determined. Amplified VH and VL cDNA sequences were used in a claimed process for the production of novel humanised, reshaped MAB 15 having humanised, reshaped VH and VL regions (see W21652 and W21651), which can be used for treating tumours, especially lung cancer, and for the manufacture of a drug related to tumours, especially lung cancer.	
CC	Sequence 140 AA;	
CC	50	

	Query	Match	Score	DB 1	Length
Best Local	Similarity	80.8%	Score 730;	DB 1;	Length 140;
Best Local	Similarity	82.1%	Pred. No. 1.19e-49;		
Matches	101;	Conservative	10;	Mismatches 9;	Indels 3; Gaps
Ddb	20	EVQVEESGGGLVKPGSQLKLSAACSGFAFSYDMMWVROTPEKKIEVAYLSRGCGSTYY	79		
Ddb	1	EVQVEESGGGLVKPGSQLKLSAACSGFAFSYDMMWVROTPEKKIEVAYLSRGCGSTYY	60		
Ddb	80	PDYVKGRFTISDRNAAKTLFLQMTSLKSEDAAMYCARHGEVVRPW--FEDYWGQSTLV	136		
Ddb	61	PDYVKGRFTISDRNAAKNTLYLQMSSLKSEDTAMYCARHSGYGSYGVLFAYWGQSTLV	120		
Ddb	137	VSA 139			
Ddb	121	VSA 123			

RESULT 7 ID R06251 standard; protein: 136 kDa.
ID R0251;
AC R0251;
DDT --10-DEC-1990--(first-entry)
Variable region of murine AHT 54 heavy chain.
Interleukin-2 receptor: IL-2: tumour necrosis factor: TNF: ss.
KW KW

Query Match 80.6%; Score 728; DB 1; Length 136;
 Best Local Similarity 88.6%; Pred. No. 1.75e-49;
 Matches 109; Conservative 3; Mismatches 5; Indels 6; Gaps 0

RESULT		8	standard; Protein: 139 AA.	
Qy	1	EVLQYSGGGGLVKPGGSLKLUSCAASEFASTYDMSHVRQTPEKRLRVEWYISGGGTTYY	6	
Db	80	PDTVKGRETISRDNAKNTLYLQMQSSLKSEDTAVYCCARR-----YGLPEAYWGQQLTVT	1	
Db	61	PDTVKGRETISRDNAKNTLYLQMQSSLKSEDTAMYTCARHSGYGSSYGVLFAYWGQQLTV	1	
Qy	134	VSA 136		
Db	121	VSA 123		
Qy	121	VSA 123		
			Location/Qualifiers	
			20..139	
			/Label= KC-4_mature_YL-chain	
			20..49	
			FT	
			FT	
			FT	

PT:	region	50.	.54
PT:	region	/label=	CDR1
PT:	PT:	55.	.68
PT:	region	/label=	FR2
PT:	PT:	69.	.85
PT:	region	/label=	CDR2
PT:	PT:	86.	.117
PT:	region	/label=	FR3
PT:	PT:	118.	.128
PT:	region	/label=	CDR3
PT:	PT:	129.	.139
PT:	region	/label=	FR4
PN:			
PN:		W09411508-A.	
PD:		26-MAY-1994.	
PD:		15-MAY-1993.	
PF:		U113116.	

PR 13-NOV-1992; US-977707.
PR 28-SEP-1993; US-128015.
PA (CANCER RES FUND CONTRA COSTA,
WPI; 94-183309/22.
DR -N-PSDB-062264-
PR Chimeric human-murine polypeptide(s),
PR fat globule antigen - for imaging, di-

PS Example 27: Page 41; 54pp; English.
 CC An initial isolation of cDNAs coding for murine anti-human breast
 CC carcinoma Mab KC-4 was performed using PCR with commercially
 CC available primers (see 062751-062758, available from NOVAGEN).
 CC Subsequent cloning using PCR primers J020, J021, J022 and J024
 CC (see 062759-062762), resulted in the isolation of the mouse Ig
 CC variable domains. The amplified cDNAs were sequenced (062763 and
 CC 062764) and amino acid sequences were deduced from them. Chimeric
 CC mouse-human antibodies were constructed using human constant
 CC regions so as to produce less immunogenic polypeptides which
 CC retained the anti-human carcinoma binding specificity of KC-4.
 SQ sequence 139 AA;

Db	20	EVQNEGGGLVPGGSKLSCANSGFAPSYYAMSIVRSPKLEWAEIISGNYAII
QY	1	EVQIVESGGGLVPGGSKLSCANSGFAPSYYAMSIWVROTSPKLEWAEIISGGTTIV
Db	80	QDTIVTGRFTIISDNAAKNTLYLEMSLRSDEAMYCARED-YGIP-A-WFAVNGQGTLYS

Qy	61	PDTVKGRFTISRDNAKNTLYQMSLKLSEDTAMYCARRHSGGSSYGVLFAYWQGQTLVT 120
Db	137	VSA 139
Qy	121	VSA 123
RESULT	9	
ID	R52791	standard; Protein: 139 AA.
AC	R52791;	
DT	24-JAN-1995	(first entry)
DE	Murine KC-4 immunoglobulin heavy chain variable region (deduced).	
KW	Immunoglobulin variable domain; primer; polymerase chain reaction;	
KW	chimeric antibody; human milk fat globule; human breast carcinoma;	
KW	murine anti-human carcinoma monoclonal antibody KC-4.	
OS	Mus musculus.	
PH	Location/Qualifiers	
FT	protein	20..139
FT	region	/label= KC-4.mature.VL-chain
FT	region	20..49
FT	region	/label= FR1
FT	region	50..54
FT	region	/label= CDRI
FT	region	55..68
FT	region	/label= FR2
FT	region	69..85
FT	region	/label= CDR2
FT	region	86..117
FT	region	/label= FR3
FT	region	118..128
FT	region	/label= CDR3
FT	region	129..139
FT	region	/label= FR4
PR	W09411509-A.	
PD	19-NOV-1998	
PD	18-JUL-1997	
PD	15-MAY-1997	
PA	(CHUS) CHUGAI SEIYAKU KK.	
PI	Ishii K, Sato K, Tunenari T;	
DR	WPI: 99-070101/06.	
PT	PT Inhibitors of binding of parathyroid hormone related peptide to its receptor - useful for, e.g. treatment of cachexia arising from cancer or other diseases.	
PS	Example 2: Page 72-73; 125PP; Japanese	
CC	The present invention describes compositions for the treatment of cachexia containing a substance which inhibits the binding of a parathyroid hormone related peptide (pTHRP) to its receptor as an active component. This substance may be an antagonist to the receptor, or an antibody (preferably monoclonal) or antibody fragment, recognising pTHRP. The antibody is preferably humanised or chimeric.	
CC	The present invention also describes a humanised antibody prepared by hybridoma 23-57-137-1 (FERM-PB5631). The composition is used for the treatment of cachexia arising in connection with diseases such as cancer, thereby improving the quality of life of the patient. The present sequence represents the mouse humanised antibody heavy chain from	
CC	CC #22-57-137-1 from the present invention.	
SQ	Sequence 118 AA;	
Query Match	79.3%	Score 716; DB 1; Length 118;
Best Local Similarity	82.9%	Pred. No. 1 74e-48;
Matches	102;	Indels 5; Gaps 7;
AC	W57576;	
DB	03-SEP-1998	(first entry)
DE	Chimeric H chain SEQ ID NO:45 for an antibody against hTHRP.	
KW	Chimeric; antibody; human parathyroid hormone related Peptide; hTHRP; mouse;	
KW	L chain; H chain; hypercalcaemia; cancer; malignant lymphoma; CDR;	
KW	hypophosphaemia; pathogen; vitamin D resistance; V region; C region;	
KW	humanised.	
KW	Synthetic.	
OS	Chimeric - Mus sp.	
OS	Chimeric - Homo sapiens.	
PN	W0913386-A1.	
PD	02-APR-1998.	
PF	24-SEP-1997; J03382.	
PR	24-JUL-1997; JP-214168.	
PR	26-SEP-1996; JP-255196.	
PA	(CHUS) CHUGAI SEIYAKU KK.	
PI	Sato K, Wakahara Y, Yabuta N;	
Qy	61	PDTVKGRFTISRDNAKNTLYQMSLKLSEDTAMYCARRHSGGSSYGVLFAYWQGQTLVT 120
Db	137	VSA 139
Qy	121	VSA 123
RESULT	11	
ID	W57576	standard; protein: 118 AA.
AC	W57576;	
DB	03-SEP-1998	(first entry)
DE	Chimeric H chain SEQ ID NO:45 for an antibody against hTHRP.	
KW	Chimeric; antibody; human parathyroid hormone related Peptide; hTHRP; mouse;	
KW	L chain; H chain; hypercalcaemia; cancer; malignant lymphoma; CDR;	
KW	hypophosphaemia; pathogen; vitamin D resistance; V region; C region;	
KW	humanised.	
KW	Synthetic.	
OS	Chimeric - Mus sp.	
OS	Chimeric - Homo sapiens.	
PN	W0913386-A1.	
PD	02-APR-1998.	
PF	24-SEP-1997; J03382.	
PR	24-JUL-1997; JP-214168.	
PR	26-SEP-1996; JP-255196.	
PA	(CHUS) CHUGAI SEIYAKU KK.	
PI	Sato K, Wakahara Y, Yabuta N;	

WPI: 98-230640/20.
New chimeric antibodies against human parathormone related peptide(s) - useful for, e.g. treatment of hypercalcaemia and other disorders caused by malignant neoplasm(s).

PT Claim 5; Page 111-112; 182pp; Japanese.

CC New antibodies have been developed which are specific for human parathormone related peptides (hPRP). The antibodies comprise chimeric L and/or H chains, where the C region is of human and L region of mouse, origin. The present sequence represents a specifically claimed region of an antibody of the invention. Host cells, transformed with vectors containing DNA encoding antibodies of the invention, can be used to produce the antibodies. The antibodies may be used to treat hypercalcaemia, especially that due to a malignancy, e.g. cancers of pancreas, lung, throat, larynx, tongue, gum, oesophagus, stomach, liver, breast, kidney, bladder, womb or prostate or malignant lymphoma. They may also be used for treatment of hypophosphaemia such as that due to pathogens or to vitamin D resistance.

CC Sequence 118 AA;

SQ Query Match 79.3%; Score 716; DB 1; Length 118;
Best Local Similarity 82.9%; Pred. No. 1.74e-48;
Matches 102; Conservative 9; Mismatches 7; Indels 5; Gaps 2;

Db 1 EVOLVESGGDLVKPGGLKLSCAAASGFTSSYGMWIROTPDKRLEWVATISGGSYTY 60
Qy 1 EVOLVESGGDLVKPGGLKLSCAAASGFAFSIYDMSWVROPEKRLWVAYISGGTY 60

Db 61 PDSVKGRTFSRDNAKNTLYLQMSLKSEDTAMYCARQT-T-MY--FAYWQGTLVT 115
Qy 61 PDTVKGRFTISRDNAKNTLYLQMSLKSEDTAMYCARHSSGSSYGVLFAYWQGTLVT 120

Db 80 PDSVKGRTFSRDNAKNTLYLQMSLKSEDTAMYCARQT-T-MY--FAYWQGTLVT 134
Qy 61 PDTVKGRFTISRDNAKNTLYLQMSLKSEDTAMYCARHSSGSSYGVLFAYWQGTLVT 120

Db 135 VSA 137
Qy 121 VSA 123

RESULT 13
ID W57592 standard; protein: 137 AA.

AC W57592; 03-SEP-1998 (first entry)

DE Chimeric antibody against hPRP H chain V region SEQ ID NO:57.

KW Chimeric; antibody; human parathormone related peptide; hPRP; mouse; L chain; H chain; hypercalcaemia; cancer; malignant lymphoma; CRR; KW hypophosphaemia; pathogen; vitamin D resistance; V region; C region; KW humanised.

OS Synthetic.

OS Chimeric - Mus sp.

OS Chimeric - Homo sapiens.

PN WO9813388-A1.

PD 02-APR-1998.

PP 24-SEP-1997; J03382.

PR 24-JUL-1997; JP-214168.

PR 26-SEP-1996; JP-255196.

PA (CHUS) CHUGAI SEIYAKU KK.

DI Sato K, Wakaha T, Yabuta N;

DR WPI; 98-230640/20.

N-PSDB; V24232.

PT New chimeric antibodies against human parathormone related peptide(s) - useful for, e.g. treatment of hypercalcaemia and other

PT disorders caused by malignant neoplasm(s)

PS Claim 52; Page 120-121; 182pp; Japanese.

CC New antibodies have been developed which are specific for human

CC -parathormone-related-peptides (hPRP) - the antibodies comprise chimeric L and/or H chains, where the C region is of human and L region of mouse, origin. The present sequence represents a specifically claimed region of an antibody of the invention. Host cells, transformed with vectors

CC containing DNA encoding antibodies of the invention, can be used to produce the antibodies. The antibodies may be used to treat

CC hypercalcaemia, especially that due to a malignancy, e.g. cancers of

CC pancreas, lung, throat, larynx, tongue, gum, oesophagus, stomach, liver, breast, kidney, bladder, womb or prostate or malignant lymphoma. They

CC may also be used for treatment of hypophosphaemia such as that due to

CC pathogens or to vitamin D resistance.

SQ Sequence 137 AA;

Query Match 79.3%; Score 716; DB 1; Length 137;

Best Local Similarity 82.9%; Pred. No. 1.74e-48;

Matches 102; Conservative 9; Mismatches 7; Indels 5; Gaps 2;

Db 20 EVOLVESGGDLVKPGGLKLSCAAASGFTSSYGMWIROTPDKRLEWVATISGGSYTY 79
Qy 1 EVOLVESGGDLVKPGGLKLSCAAASGFAFSIYDMSWVROPEKRLWVAYISGGTY 60

Db 80 PDSVKGRTFSRDNAKNTLYLQMSLKSEDTAMYCARQT-T-MY--FAYWQGTLVT 134
Qy 61 PDTVKGRFTISRDNAKNTLYLQMSLKSEDTAMYCARHSSGSSYGVLFAYWQGTLVT 120

Db 135 VSA 137

CC #23-57-137-1 from the present invention.

SQ Sequence 137 AA;

Query Match 79.3%; Score 716; DB 1; Length 137;

Best Local Similarity 82.9%; Pred. No. 1.74e-48;

Matches 102; Conservative 9; Mismatches 7; Indels 5; Gaps 2;

Db 20 EVOLVESGGDLVKPGGLKLSCAAASGFTSSYGMWIROTPDKRLEWVATISGGSYTY 79
Qy 1 EVOLVESGGDLVKPGGLKLSCAAASGFAFSIYDMSWVROPEKRLWVAYISGGTY 60

Db 80 PDSVKGRTFSRDNAKNTLYLQMSLKSEDTAMYCARQT-T-MY--FAYWQGTLVT 134
Qy 61 PDTVKGRFTISRDNAKNTLYLQMSLKSEDTAMYCARHSSGSSYGVLFAYWQGTLVT 120

Db 135 VSA 137

Qy 121 VSA 123

Qy	121 VSA 123	Db	80 PDTVKGRFTISRDNAKNTLYQMSLSSKSEDTAMYHCAAR-GGVRGGY--EDVWGAGTVT 135
RESULT	14	Qy	61 PDTVKGRFTISRDNAKNTLYQMSLSSKSEDTAMYHCAAR-HSGYGSYCVLFAYWQGFLVT 120
ID	W11917; standard; Protein; 247 AA.	Db	136 VSS 138
AC	W11917;	Qy	121 VSA 123
DT	24-JUN-1997 (first entry)		
DE	Interleukin-1 heavy chain.		
KW	Interleukin-1 beta; IFN-1 beta; recombinant antibody; humanised antibody; chimeric antibody; antibody engineering; monoclonal antibody; MAB; SK48-E26; inflammation; therapy.		
OS	Homo sapiens.		
FF			
Key peptide			
FT			
FT	1.19		
FT	/label= Sig_peptide		
FT	20.49		
FT	/label= FR1		
FT			
FT	/note= "framework region 1"		
FT	50.54		
FT	/label= CDR1		
FT			
FT	/note= "complementarity determining region 1"		
FT	(Claim 10, page 48).		
FT	55.68		
FT	/label= FR2		
FT			
FT	/note= "framework region 2"		
FT	69.85		
FT	/label= CDR2		
FT			
FT	/note= "complementarity determining region 2"		
FT	(Claim 10, page 48).		
FT	86.117		
FT	/label= FR3		
FT			
FT	/note= "framework region 3"		
FT	118.127		
FT	/label= CDR3		
FT			
FT	/note= "complementarity determining region 3"		
FT	Claim 10, page 48).		
FT	128.138		
FT	/label= FR4		
FT			
FT	/note= "framework region 4"		
FT	139.247		
FT	/label= Constant_region		
FT			
FT	W09501997-A1.		
FT	PD 19-JAN-1995.		
FT	07-JUL-1994; US-07659.		
FT	09-JUL-1993; US-090534.		
FT	04-MAR-1994; US-206190.		
PA	(SMIK) SMITHKLINE BEECHAM CORP.		
PI	Gross MS, Hurie MR, Jackson JR, Jonak ZL, Theisen TW;		
PI	Young PR;		
WPI	95-066868/09.		
DR	N-PSDB; T51436.		
PT	Recombinant and humanised chimeric antibodies against human		
PT	interleukin-1 beta - for preventing and treating		
PT	interleukin-mediated inflammatory disorders		
PS	Claim 5; Page 36-37; 62pp; English.		
CC	Amino acid sequences of the heavy chain (W11917) and light chain		
CC	(W11918) of anti-human interleukin 1 beta (IL-1 beta) murine		
CC	monoclonal antibody (MAB) SK48-E26 were deduced from nucleic acids		
CC	(T51436-37) derived from hybridoma SK48-E26. The heavy and light		
CC	chains, esp. the complementarity determining region sequences,		
CC	can be utilised in novel recombinant chimeric and humanised		
CC	antibodies (see also W11919-20) useful for the treatment and		
CC	prevention of IL-1 mediated inflammatory disorders.		
SQ	Sequence 247 AA;		
Query Match	79.3%	Score 716; DB 1; Length 247;	
Best Local Similarity	84.8%	Pred. No. 1.74e-48;	
Matches	104; Conservative	5; Mismatches 10; Indels 4; Gaps 2;	
Db	20 EVH1VEGGGLVPGGGSLKLSCAAGFAFSSYDMSWROTPEKRLDWWAYISSGGGCTY	1	
Qy	1 EVQLVESGGGLVPGGGSLKLSCAAGFAFSSYDMSWROTPEKRLDWWAYISSGGGCTY	1	

Search completed: Mon Oct 2 09:33:25 2000
Job time : 14 secs.

Sat Oct 7 12:43:25 2000

US-09-381-49

ID R84553 standard; Protein; 131 AA.
IC R84553;
JT 02-FEB-1996 (first entry)
DE Mab SCH94.03 light chain.
KW Monoclonal antibody; Mab; SCH94.03; hybridoma; central nervous system;
KW CNS; demyelination; multiple sclerosis; neural disease; therapeutic.
DS Mus sp.
FH Key Location/Qualifiers
FT peptide 1. .20
FT /label- Leader_peptide
FT region 44. .54
FT /label- CDR1
FT region 70. .76
FT /label- CDR2
FT region 109. .117
FT /label- CDR3
FT region 116. .128
FT /label- Joining_region
FT region 129. .131
FT /label- C-kappa_region
PN WO9530004-A1.
PD 09-NOV-1995.
PF 27-APR-1995; U05262.
PR 29-APR-1994; US-236520.
PA (MAYO-) MAYO FOUNDATION.
PI Miller DJ, Rodriguez M;
DR WPI; 95-393077/50.
DR N-PSDB; T05311.
PT Monoclonal antibodies which stimulate central nervous system
PT re-myelination - are produced by hybridoma ATCC CRL 11627, for
PT treating multiple sclerosis, and viral or post-neuronal diseases of
PT the CNS.
PS Disclosure: Page 36-37; 63pp; English.
CC Hybridoma ATCC CRL 11627 was obtd. from a SJL/J mouse injected with
CC spinal cord homogenate from a mammal uninfected with any
CC demyelinating disease. The hybridoma produced a monoclonal antibody
CC (SCH94.03) useful in promoting CNS remyelination. The SCH94.03
CC light chain amino acid sequence is given in R84553.
SO Sequence 131 AA;

```
Query Match      97.0%;  Score 734;  DB 1;  Length 131;
Best Local Similarity 97.2%;  Pred. No. 9.32e-50;
Matches 104;  Conservative 2;  Mismatches 1;  Indels 0;  Gaps 0;
```

DB 21 DIQMTQTTSSLASLGDRVТИSCRASQDISNYLNWYQQKPDGTVKLLIYYTSRLHSGVPS 80

Ov 1 D10MTOTSSL-SASLGDRVT1SCRASQDISNYLNHYQQKPDGTVKLL1YYTSILHSGVPS 60

ПРИЛОЖЕНИЕ 1. ПРИМЕРЫ ПОДСЧЕТЫ ВЫПРЯМЛЕНЫХ КОМПЛЕКСОВ

Qy 61 KFSGSGSGTDYSLTISNLEQEDFATYFCQQGNTLPWTGGGTLEIK 107